#8

SEQUENCE LISTING

<110> Miki Dr., Brian L

Gijzen Dr., Matk

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Boutilier Dr., Kim

Hu., Ming

Bowman, LuAnn

Batchelor, Anthea

<120> Seed-Coat Promoters, Genes and Gene Poducts

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20 25 30

Pro Gln Pro Ser His Val Thr Ala Leu Ile Thr Arg Pro Ser Cys Pro

35 40 45

Asp Leu Ser Ile Cys Leu Asn Ile Leu Gly Gly Ser Leu Gly Thr Val

50 55 60

Asp Asp Cys Cys Ala Leu Ile Gly Gly Leu Gly Asp Ile Glu Ala Ile

65 70 75 80

Val Cys Leu Cys Ile Gln Leu Arg Ala Leu Gly Ile Leu Asn Leu Asn

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Arg Asn Leu Gln Leu Ile Leu Asn Ser Cys Gly Arg Ser Tyr Pro Ser

100 105 110

Asn Ala Thr Cys Pro Arg Thr

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Ser Ile Phe Glu Tyr Ala Gly Asn Asp Asp Gln Ser Glu Ser Arg Ser

35 40 45

Ile Leu Gly Tyr Ala Gly Tyr Asn Gln Asp Glu Asp Asp Val Ser Lys

50 55 60

His Asn Ile Gln Ile Phe Asn Arg Leu Phe Phe Leu Glu Glu Asp Leu

65 70 75 80

Arg Ala Gly Lys Ile Phe Asn Met Lys Phe Val Asn Asn Thr Lys Ala

85 90 95

Thr Val Pro Leu Leu Pro Arg Gln Ile Ser Lys Gln Ile Pro Phe Ser

100 105 110

Glu Asp Lys Lys Lys Gln Val Leu Ala Met Leu Gly Val Glu Ala Asn

115 120 125

Ser Ser Asn Ala Lys Ile Ile Ala Glu Thr Ile Gly Leu Cys Gln Glu

130 135 140

Pro Ala Thr Glu Gly Glu Arg Lys His Cys Ala Thr Ser Leu Glu Ser

145 150 155 160

Met Val Asp Phe Val Val Ser Ala Leu Gly Lys Asn Val Gly Ala Phe

165 170 175

Ser Thr Glu Lys Glu Arg Glu Thr Glu Ser Gly Lys Phe Val Val Val

180 185 190

Lys Asn Gly Val Arg Lys Leu Gly Asp Asp Lys Val Ile Ala Cys His

195 200 205

Pro Met Ser Tyr Pro Tyr Val Val Phe Gly Cys His Leu Val Pro Arg

210 215 220

Ser Ser Gly Tyr Leu Val Arg Leu Lys Gly Glu Asp Gly Val Arg Val

225 230 235 240

Lys Ala Val Val Ala Cys His Arg Asp Thr Ser Lys Trp Asp His Asn

245 250 255

His Gly Ala Phe Lys Val Leu Asn Leu Lys Pro Gly Asn Gly Thr Val

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Cys His Val Phe Thr Glu Gly Asn Leu Leu Trp Leu Pro Asn

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<210> 12

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<212> PRT

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Met Lys Gly Asn Asn Thr Leu Leu Leu His Leu Phe Tyr Thr Thr Leu

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Phe Leu Phe Leu Val Val Ser Ser Ser Ser Ser Thr Gly Asn Glu Ser

20 25 30

Asn Asp Asp Thr Asn Ser Lys Glu Val Tyr Ile Val Tyr Met Gly Ala

35 40 45

Ala Asp Ser Thr Lys Ala Ser Leu Lys Asn Glu His Ala Gln Ile Leu

50 55 60

Asn Ser Val Leu Arg Arg Asn Glu Asn Ala Leu Val Arg Asn Tyr Lys

65 70 75 80

His Gly Phe Ser Gly Phe Ala Ala Arg Leu Ser Lys Glu Glu Ala Asn

85 90 95

Ser Ile Ala Gln Lys Pro Gly Val Val Ser Val Phe Pro Asp Pro Ile

100 105 110

Leu Lys Leu His Thr Thr Arg Ser Trp Asp Phe Leu Lys Ser Gln Thr

115 120 125

Arg Val Asn Ile Asp Thr Lys Pro Asn Thr Leu Ser Gly Ser Ser Phe

130 135 140

Ser Ser Ser Asp Val Ile Leu Gly Val Leu Asp Thr Gly Ile Trp Pro

145 150 155 160

Glu Ala Ala Ser Phe Ser Asp Lys Gly Phe Gly Pro Val Pro Ser Arg

165 170 175

Trp Lys Gly Thr Cys Met Thr Ser Lys Asp Phe Asn Ser Ser Cys Cys

180 185 190

Asn Arg Lys Ile Ile Gly Ala Arg Phe Tyr Pro Asn Pro Glu Glu Lys

195 200 205

Thr Ala Arg Asp Phe Asn Gly His Gly Thr His Val Ser Ser Thr Ala

210 215 220

Val Gly Val Pro Val Ser Gly Ala Ser Phe Tyr Gly Leu Ala Ala Gly

225 230 235 240

Thr Ala Arg Gly Gly Ser Pro Glu Ser Arg Leu Ala Val Tyr Lys Val

245 250 255

Cys Gly Ala Phe Gly Ser Cys Pro Gly Ser Ala Ile Leu Ala Gly Phe

260 265 270

Asp Asp Ala Ile His Asp Gly Val Asp Ile Leu Ser Leu Ser Leu Gly

275 280 285

Gly Phe Gly Gly Thr Lys Thr Asp Leu Thr Thr Asp Pro Ile Ala Ile

290 295 300

Gly Ala Phe His Ser Val Gln Arg Gly Ile Leu Val Val Cys Ala Ala

305 310 315 320

Gly Asn Asp Gly Glu Pro Phe Thr Val Leu Asn Asp Ala Pro Trp Ile

325 330 335

Leu Thr Val Ala Ala Ser Thr Ile Asp Arg Asp Leu Gln Ser Asp Val

340 345 350

Val Leu Gly Asn Asn Gln Val Val Lys Gly Arg Ala Ile Asn Phe Ser

355 360 365

Pro Leu Leu Asn Ser Pro Asp Tyr Pro Met Ile Tyr Ala Glu Ser Ala

370 375 380

Ala Arg Ala Asn Ile Ser Asn Ile Thr Asp Ala Arg Gln Cys His Pro

385 390 395 400

Asp Ser Leu Asp Pro Lys Lys Val Ile Gly Lys Ile Val Val Cys Asp

405 410 415

Gly Lys Asn Asp Ile Tyr Tyr Ser Thr Asp Glu Lys Ile Val Ile Val

420 425 430

Lys Ala Leu Gly Gly Ile Gly Leu Val His Ile Thr Asp Gln Ser Gly

435 440 445

Ser Val Ala Phe Tyr Tyr Val Asp Phe Pro Val Thr Glu Val Lys Ser

450 455 460

Lys His Gly Asp Ala Ile Leu Gln Tyr Ile Asn Ser Thr Ser His Pro

465 470 475 480

Val Gly Thr Ile Leu Ala Thr Val Thr Ile Pro Asp Tyr Lys Pro Ala

485 490 495

Pro Arg Val Gly Tyr Phe Ser Ser Arg Gly Pro Ser Leu Ile Thr Ser

500 505 510

Asn Val Leu Lys Pro Asp Ile Ala Ala Pro Gly Val Asn Ile Leu Ala

515 520 525

Ala Trp Phe Gly Asn Asp Thr Ser Glu Val Pro Lys Gly Arg Lys Pro

530 535 540

Ser Leu Tyr Arg Ile Leu Ser Gly Thr Ser Met Ala Thr Pro His Val

545 550 555 560

Ser Gly Leu Ala Cys Ser Val Lys Arg Lys Asn Pro Thr Trp Ser Ala

565 570 575

Ser Ala Ile Lys Ser Ala Ile Met Thr Ser Ala Ile Gln Asn Asp Asn

580 585 590

Leu Lys Gly Pro Ile Thr Thr Asp Ser Gly Leu Ile Ala Thr Pro Tyr

595 600 605

Asp Tyr Gly Ala Gly Ala Ile Thr Thr Ser Glu Pro Leu Gln Pro Gly

610 615 620

Leu Val Tyr Glu Thr Asn Asn Val Asp Tyr Leu Asn Tyr Leu Cys Tyr

625 630 635 640

Asn Gly Leu Asn Ile Thr Met Ile Lys Val Ile Ser Gly Thr Val Pro

645 650 655

Glu Asn Phe Asn Cys Pro Lys Asp Ser Ser Ser Asp Leu Ile Ser Ser

660 665 670

Ile Asn Tyr Pro Ser Ile Ala Val Asn Phe Thr Gly Lys Ala Asp Ala

675 680 685

Val Val Ser Arg Thr Val Thr Asn Val Asp Glu Glu Asp Glu Thr Val

690 695 700

Tyr Phe Pro Val Val Glu Ala Pro Ser Glu Val Ile Val Thr Leu Phe

705 710 715 720

Pro Tyr Asn Leu Glu Phe Thr Thr Ser Ile Lys Lys Gln Ser Tyr Asn

725 730 735

Ile Thr Phe Arg Pro Lys Thr Ser Leu Lys Lys Asp Leu Phe Gly Ser

740 745 750

Ile Thr Trp Ser Asn Asp Lys Tyr Met Val Arg Ile Pro Phe Val Leu

755 760 765

Thr Lys

770